Figure S3. Characterization of the Cbk1 hydrophobic motif and its binding to the Cbk1–Mob2 complex

A

<table>
<thead>
<tr>
<th></th>
<th>Direct $K_d$ (μM)</th>
<th>Competitive $K_d$ with HM-P (μM)</th>
<th>Competitive $K_d$ with pepSSD1 (μM)</th>
</tr>
</thead>
<tbody>
<tr>
<td>HM peptide</td>
<td>33</td>
<td>ND</td>
<td>ND</td>
</tr>
<tr>
<td>HM-P peptide</td>
<td>6</td>
<td>45</td>
<td>No competitive binding</td>
</tr>
</tbody>
</table>

B

Direct binding of pepHM

Kd = 32.6 ± 7.7 μM

Direct binding of pepHM-P

Kd = 6.3 ± 0.6 μM

C

Competitive binding of pepHM-P

Kd = 45 ± 18 μM

D

Competitition of pepHM-P and pepSSD1

No competitive binding